



1600
1642
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RAW SEQUENCE LISTING

DATE: 03/29/2002

PATENT APPLICATION: US/09/960,631A

TIME: 13:28:15

Input Set : A:\266_171.ST25.txt

Output Set: N:\CRF3\03292002\I960631A.raw

3 <110> APPLICANT: MIROCHNITCHENKO, Oleg
 4 WEI, Jiang
 5 INOUE, Masayori
 7 <120> TITLE OF INVENTION: SOLUBLE ISCHEMIA ACTIVATED PROTEIN
 9 <130> FILE REFERENCE: 266/171
 11 <140> CURRENT APPLICATION NUMBER: US 09/960,631A
 12 <141> CURRENT FILING DATE: 2001-09-20
 14 <150> PRIOR APPLICATION NUMBER: US 60/233,819
 15 <151> PRIOR FILING DATE: 2000-09-20
 17 <160> NUMBER OF SEQ ID NOS: 8
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 840
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Homo sapiens
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 31 gcggcccccg gcgcccggct gttgcggctc ccggggagcg gggccgtgca ggccgcgagc 180
 33 ccggagcgcg ccggttgga cagggcgctg cgggccgcgg tggccgagct gcgcgcgggc 240
 35 gccgtggtgg ccgtccccc cgtatcgctg tacggcctgg cctgcgcggc gagctgctcg 300
 37 gcggctctgc gcgctgtgta ccgcctcaag ggtcgagcg aggccaaagg tctggccgta 360
 39 tgccctggcc gcgtggccga cgtctacaga tactgccgtg tgagagtacc tgaggggctc 420
 41 ctgaaagacc tactgccagg accagtgaac ctggtgatgg aacgctcgga ggagctcaac 480
 43 aaggacctaa accctttttac gcctcttgta ggcattcgga ttcttgatca tgcttttatg 540
 45 caagacttgg ctcatagtt tgaggggtccg cttgctctca ctagtccaa cctcagctcc 600
 47 caggccagtt ctctgaatgt cgaggagttc caggatctct ggccctcagtt gtccttggtt 660
 49 attgatgggg gacaaattgg ggatggccag agccccgagt gtcgccttgg ctcaactgtg 720
 51 gttgatttgt ctgtgcccgg aaagtttggt atcattcgtc caggctgtgc cctggaaagt 780
 53 actacagcca tctccaaca gaagtacgga ctgctcccct cacatgcgtc ctacctgtga 840
 56 <210> SEQ ID NO: 2
 57 <211> LENGTH: 279
 58 <212> TYPE: PRT
 59 <213> ORGANISM: Homo sapiens
 61 <400> SEQUENCE: 2
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 64 1 5 10 15
 67 Ser Val Gly Leu Ser Glu Gly Pro Ala Gly Ser Arg Ser Gly Arg Leu
 68 20 25 30
 71 Phe Arg Pro Pro Ser Pro Ala Pro Ala Ala Pro Gly Ala Arg Leu Leu
 72 35 40 45
 75 Arg Leu Pro Gly Ser Gly Ala Val Gln Ala Ala Ser Pro Glu Arg Ala
 76 50 55 60

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79 Gly Tyr Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala Gly
80 65          70          75          80
83 Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys Ala
84          85          90          95
87 Ala Ser Cys Ser Ala Ala Leu Arg Ala Val Tyr Arg Leu Lys Gly Arg
88          100          105          110
91 Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp Val
92          115          120          125
95 Tyr Arg Tyr Cys Arg Val Arg Val Pro Glu Gly Leu Leu Lys Asp Leu
96          130          135          140
99 Leu Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu Asn
100 145          150          155          160
103 Lys Asp Leu Asn Pro Phe Thr Pro Leu Val Gly Ile Arg Ile Pro Asp
104          165          170          175
107 His Ala Phe Met Gln Asp Leu Ala Gln Met Phe Glu Gly Pro Leu Ala
108          180          185          190
111 Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Asn Val Glu
112          195          200          205
115 Glu Phe Gln Asp Leu Tyr Pro Gln Leu Ser Leu Val Ile Asp Gly Gly
116          210          215          220
119 Gln Ile Gly Asp Gly Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr Val
120 225          230          235          240
123 Val Asp Leu Ser Val Pro Gly Lys Phe Gly Ile Ile Arg Pro Gly Cys
124          245          250          255
127 Ala Leu Glu Ser Thr Thr Ala Ile Leu Gln Gln Lys Tyr Gly Leu Leu
128          260          265          270
131 Pro Ser His Ala Ser Tyr Leu
132          275
135 <210> SEQ ID NO: 3
136 <211> LENGTH: 1387
137 <212> TYPE: DNA
138 <213> ORGANISM: Homo sapiens
140 <220> FEATURE:
141 <221> NAME/KEY: misc_feature
142 <222> LOCATION: (1)..(1387)
143 <223> OTHER INFORMATION: The letter "n" stands for a substitution base.
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149 cttgtaggca ttcggattcc tgatcatgct tttatgcaag acttggtcca gatgtttgag      120
151 ggtccgcttg ctctcactag tgccaacctc agctcccagg ccagttctct gaatgtcgag      180
153 gagttccagg atctctggcc tcagttgtcc ttggttattg atgggggaca aattggggat      240
155 ggccagagcc ccgagtgtcg ccttggtcca actgtggttg atttgtctgt gcccggaaag      300
157 tttggcatca ttggtccagg gtgtgcctgg gaaagtacta cagccatcct ccaacagaag      360
159 tacggactgc tcccctcaca tgcgtcctac ctgtgaaact ctgggaagca ggaaggccca      420
161 agacctggtg ctggatacta tgtgtctgtc cactgacgac tgtcaaggcc tcatttgcag      480
163 aggccaccgg agctagggca ctagcctgac ttttaaggca gtgtgtcttt ctgagcactg      540
165 tagaccaagc ccttggaagt gctgggttag ccttgcaact ggggaaagga tgtattttatt      600
167 tgtattttca tatatcagcc aaaagctgaa tggaaaagtt aagaacattc ctaggtggcc      660
169 ttattctaata aggtttcttc tgtctgtttt gtttttcaat tgaaaagtaa ttaaataaca      720

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Output Set: N:\CRF3\03292002\I960631A.raw

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W--> 171 gatttagaat ctagtgagag cntcctctct gggggtggtg gcatttaagg ttcaaccan 780
W--> 173 ccnagaagtg ctgcgctgtt taaaaagtct caggtggctg cgtgtggtgg ctcatgctg 840
175 taatcccaac attctgggag gcccaggcgg gagaactgct tgagcccagg agttcagaat 900
177 cagcctgggc aacatagcaa tactccgtct cataaaaatt aataaataaa aagtctcagg 960
179 tgaccaaaagg ctccctgaagc tagaaccagg tttggataaa gattgaagag ccacaggcca 1020
181 ctcttccctc tgagccattg ggcctagtgg tgtcatgtat tgtaattgct cgcagggaga 1080
183 gcagtctttt tgggtgaata gtgggatgtc tgcttagttg gcaggggttc agtccaaatg 1140
W--> 185 gaagaatatt gggaaataaa cctccnctat cctttatagc cagggaacttt tttcttattt 1200
187 attcataaaa taaattatag ttaattatac ccataacacc tttatttaaa tccagtgttc 1260
189 tccgcagcct tttgtctatt tataatgtgta ccaagtgtta aacataatta ttattgggca 1320
W--> 191 tttgaacntg tttttcntta naganatnct gnnattaaac atatttgtna atggnaaaaa 1380
193 aaaaaaa 1387
196 <210> SEQ ID NO: 4
197 <211> LENGTH: 930
198 <212> TYPE: DNA
199 <213> ORGANISM: Mus musculus
201 <400> SEQUENCE: 4
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204 agcgacgggc cggctagtgc tggccgccc tgcgcctcc tactccctcc tgagcccgtc 120
206 ccggcgctgc cgggggccc gctgctgcgg ctcccgagaga gcgagcccgt ggaagccgcg 180
208 agccccgagc gcgcgggctg gaccgagcgc ctgcccggcc cgcgtggccga gctgcgcgcc 240
210 ggcgcgctgg tggcggtccc gaccgacacg ctctacggcc tggcctgctc ggcgagcagc 300
212 tcggcggccc tgagttgcgt gtaccgcctc aaaggccgca gcgaggccaa gccgtggcc 360
214 gtgtgcctgg gccgcgtggc cgacgtctac aggtactgtc aggtgagagt acctagggag 420
216 ctccctggaag acctgttccc aggccctgtg accctggtga tggagcgctc cgaggagctc 480
218 aacaaagacc tgaacccctt tactcgtctt gttggcatcc ggattcctga ccatgccttc 540
220 atgtctgact tggccagat gtttggggga ccacttgac tccactagtgc caacctcagc 600
222 tcccaggcca gttctctgag tgttgaggag ttccaagacc tctggcctca tttgtccctt 660
224 gtcattgatg gggggccaat tggggatagt cagagccctg agtgctgcct cggctctact 720
226 gtggttgact tatctgtgcc tggaaagtgt ggcattattc gcccaggctg tgccctggaa 780
228 aacactacat cgatcctcca gcagaaatat gggctgctcc ctccacaggg gtccgtttca 840
230 tgaaacttgg gaggacccaa ggacatgctg gatactatgt gtctgctact ggatgcaaag 900
232 cctcattgcc tgaggttcct acatctatag 930
235 <210> SEQ ID NO: 5
236 <211> LENGTH: 280
237 <212> TYPE: PRT
238 <213> ORGANISM: Mus musculus
240 <400> SEQUENCE: 5
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243 1 5 10 15
246 Gly Met Gly Leu Ser Asp Gly Pro Ala Ser Ser Gly Arg Gly Cys Arg
247 20 25 30
250 Leu Leu Leu Pro Pro Glu Pro Ala Pro Ala Leu Pro Gly Ala Arg Leu
251 35 40 45
254 Leu Arg Leu Pro Glu Ser Glu Pro Val Glu Ala Ala Ser Pro Glu Arg
255 50 55 60
258 Ala Gly Tyr Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala
259 65 70 75 80
262 Gly Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys

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263          85          90          95
266 Ser Ala Ser Ser Ser Ala Ala Leu Ser Cys Val Tyr Arg Leu Lys Gly
267          100          105          110
270 Arg Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp
271          115          120          125
274 Val Tyr Arg Tyr Cys Gln Val Arg Val Pro Arg Glu Leu Leu Glu Asp
275          130          135          140
278 Leu Phe Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu
279 145          150          155          160
282 Asn Lys Asp Leu Asn Pro Phe Thr Arg Leu Val Gly Ile Arg Ile Pro
283          165          170          175
286 Asp His Ala Phe Met Leu Asp Leu Ala Gln Met Phe Gly Gly Pro Leu
287          180          185          190
290 Ala Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Ser Val
291          195          200          205
294 Glu Glu Phe Gln Asp Leu Tyr Pro His Leu Ser Leu Val Ile Asp Gly
295          210          215          220
298 Gly Pro Ile Gly Asp Ser Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr
299 225          230          235          240
302 Val Val Asp Leu Ser Val Pro Gly Lys Phe Gly Ile Ile Arg Pro Gly
303          245          250          255
306 Cys Ala Leu Glu Asn Thr Thr Ser Ile Leu Gln Gln Lys Tyr Gly Leu
307          260          265          270
310 Leu Pro Ser Gln Gly Ser Cys Ser
311          275          280

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314 <210> SEQ ID NO: 6

315 <211> LENGTH: 702

316 <212> TYPE: DNA

317 <213> ORGANISM: Bos taurus

319 <400> SEQUENCE: 6

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324 ccgctgtggc gacgtctaca ggtactgcca cgtgagagta cctgaggggc tcctgaagga      180
326 cctgttgcca ggaccagtga ccctggtgat ggaacgctca gaggagctca acaaggacct      240
328 gaatcctttc actcctcttg taggcattccg gattcctgac caccgcttca tgcaggactt      300
330 ggtccagatg tttggggggc cactcgctct caccagtgcc aacctcagct ccagtcacag      360
332 ctctctgaat gttgaggaat tccaggacct gtggcctcac ttgtccctga tcattggtgg      420
334 gggaccaatt ggggacggcc agagcccaga gtgtcgacta ggctcaactg tggttgactt      480
336 gtctgtgcct ggaaagtittg gcatcattcg tcctggttgt gcccttgaaa gtacttcagc      540
338 catcctccag gagtatgggc tgctcccttc acatggatcc tgctggtgac actctggagg      600
340 aggggaaggcc caagggtctg tgctggacac tatgtgtccg actgctggtg gttggcaagg      660
342 cctcatttgc agaggctgct agggctacag tgtagtagt ct                          702

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345 <210> SEQ ID NO: 7

346 <211> LENGTH: 126

347 <212> TYPE: PRT

348 <213> ORGANISM: Bos taurus

350 <400> SEQUENCE: 7

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352 Met Glu Arg Ser Glu Glu Leu Asn Lys Asp Leu Asn Pro Phe Thr Pro
353 1          5          10          15

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Output Set: N:\CRF3\03292002\I960631A.raw

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356 Leu Val Gly Ile Arg Ile Pro Asp His Ala Phe Met Gln Asp Leu Val
357          20          25          30
360 Gln Met Phe Gly Gly Pro Leu Ala Leu Thr Ser Ala Asn Leu Ser Ser
361          35          40          45
364 Gln Ser Ser Ser Leu Asn Val Glu Glu Phe Gln Asp Leu Trp Pro His
365          50          55          60
368 Leu Ser Leu Ile Ile Gly Gly Gly Pro Ile Gly Asp Gly Gln Ser Pro
369 65          70          75          80
372 Glu Cys Arg Leu Gly Ser Thr Val Val Asp Leu Ser Val Pro Gly Lys
373          85          90          95
376 Phe Gly Ile Ile Arg Pro Gly Cys Ala Leu Glu Ser Thr Ser Ala Ile
377          100         105         110
380 Leu Gln Glu Tyr Gly Leu Leu Pro Ser His Gly Ser Cys Trp
381          115         120         125
384 <210> SEQ ID NO: 8
385 <211> LENGTH: 841
386 <212> TYPE: DNA
387 <213> ORGANISM: Rattus novartis
389 <220> FEATURE:
390 <221> NAME/KEY: misc_feature
391 <222> LOCATION: (491)..(491)
392 <223> OTHER INFORMATION: The letter "z" stands for sequence hybridizing.
395 <400> SEQUENCE: 8
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400 aaatatgggc tgctcccttc acaggggtcc tgttcatgaa acttgggagg acctcaagaac      180
402 catgctggat actatgtgtc tactacaggt tggcaaagcc tcattggctg aggttctctg      240
404 agctacatct gtagcctagc tttttaggca gtgtccttgg ctctgaatcc tgtaggccag      300
406 ccagaagctt cgggttgagc cttgcaccca ggggaagggt atatttactc tgtagattca      360
408 tgtgtcaacc cagaatggag ggaagaacat tcttagagtg accttattat ttttaagtgc      420
410 cctctcaccc caaccctgcc tataagttaa gtaacttgac tgcagaatta gaatgcatta      480
412 agagctgctt actggtgaac agtgaaatgt ggtttaaaac cagccagaag cactaatgca      540
414 gtctagaagt ctcaggacca atgcagcaaa gtctaggagc cctggccaga gctttctggg      600
416 tacaggagag tggtcatttg gagaaaatta ttctaggagt tccaaatgaa ataatttga      660
418 aaaataaaat cttgactgtt ttcagccagt gactttctta tttattggta tagttctctg      720
420 ttttaatttat ttaactcaga agtcattctt gtccatattg ctacctggta tttacataat      780
422 tatttttaag tatttgaact gtatttcttt attaaatatt tcttctacaa aaaaaaaaaa      840
424 a

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VERIFICATION SUMMARY

DATE: 03/29/2002

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Input Set : A:\266_171.ST25.txt

Output Set: N:\CRF3\03292002\I960631A.raw

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L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3